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DATE: Monday, May 09, 2005

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<input type="checkbox"/>	L1	dam same (dna or cdna or antisense or gene or operon or coding or encoding or nucleotide or nucleic or nuclear or polynucleotide or poly-nucleotide or rna or mrna)	1960
<input type="checkbox"/>	L2	dam.clm. same (dna or cdna or antisense or gene or operon or coding or encoding or nucleotide or nucleic or nuclear or polynucleotide or poly-nucleotide or rna or mrna).clm.	73
<input type="checkbox"/>	L3	L2 and (method or process).clm.	54
<input type="checkbox"/>	L4	mahan.in.	648
<input type="checkbox"/>	L5	L4 and salmonell\$	19
<input type="checkbox"/>	L6	l5 and (methylase or methyl-ase or methyltransferase or methyl-transferase or dam or dam-methylase or dammethylase)	13

END OF SEARCH HISTORY

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L3: Entry 15 of 54

File: PGPB

Jul 3, 2003

DOCUMENT-IDENTIFIER: US 20030124725 A1

TITLE: Mutagenesis method

CLAIMS:

1. A method for mutagenesis of a gene, which comprises introducing much more point mutations into one strand of double-stranded genomic DNA of cell or organism individual than into another strand.

2. The method according to claim 1, wherein the point mutations are randomly introduced into four kinds of bases.

3. The method according to claim 1 or 2, wherein the cell or the organism individual is mutant cell strain or mutant organism individual having mutator gene in a mutation repair gene group.

4. The method mutation according to claim 3, wherein the mutator gene is one or more mutator genes selected from a group consisting of dnaQ, dnaE, mutL, mutS, mutH, uvrD and dam.

5. The method according to claim 3 or 4, wherein the mutator gene is a gene which causes a defect of mutation repair mechanism under a certain condition.

6. The method according to claim 5, wherein the condition for the defect of the mutation repair mechanism is a certain temperature.

7. The method according to claim 5 or 6, wherein a step of introduction of mutation into genomic DNA under a certain condition and a step of selection of mutant under a selection load condition without introduction of mutation are repeated.

8. The method according to claim 7, wherein the step of introduction of mutation at the second time and thereafter are carried out under the same selection load as that in the step of mutant selection immediately therebefore.

9. A mutant of cell or organism individual where mutation is introduced into genomic DNA by any of the methods of claims 1 to 8.

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Jul 4, 2002

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CROSS-REFERENCE

[0001] This patent application is a continuation-in-part of U.S. patent application Ser. No. 09/612,116 filed Jul. 7, 2000 which is a continuation-in-part of U.S. patent application Ser. No. 09/495,614, filed Feb. 1, 2000, which claims the priority benefit of U.S. patent application Ser. Nos. 09/241,951, filed Feb. 2, 1999, converted to U.S. Provisional Ser. No. 60/183,043, and 09/305,603, filed May 5, 1999, converted to U.S. Provisional Ser. No. 60/198,250, all of which are incorporated by reference in their entirety and to which applications is claimed priority.

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L3: Entry 18 of 54

File: PGPB

Jul 4, 2002

PGPUB-DOCUMENT-NUMBER: 20020086332

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020086332 A1

TITLE: Method of reducing bacterial proliferation

PUBLICATION-DATE: July 4, 2002

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY	RULE-47
Mahan, Michael J.	Santa Barbara	CA	US	
Heithoff, Douglas M.	Goleta	CA	US	
Low, David A.	Goleta	CA	US	
Sinsheimer, Robert L.	Santa Barbara	CA	US	

APPL-NO: 09/ 928227 [\[PALM\]](#)

DATE FILED: August 9, 2001

RELATED-US-APPL-DATA:

Application 09/928227 is a continuation-in-part-of US application 09/612116, filed July 7, 2000, PENDING

Application 09/612116 is a continuation-in-part-of US application 09/495614, filed February 1, 2000, PENDING

Application is a non-provisional-of-provisional application 60/183043, filed February 2, 1999,

Application is a non-provisional-of-provisional application 60/198250, filed May 5, 1999,

INT-CL: [07] [G01](#) [N](#) [33/53](#)

US-CL-PUBLISHED: 435/7.1

US-CL-CURRENT: [435/7.1](#)

REPRESENTATIVE-FIGURES: NONE

ABSTRACT:

Bacteria and in particular pathogenic bacteria are treated in a manner which alters the bacteria's native level or activity of DNA methyltransferase (Dam). The alteration results in a change in the bacteria's native level of methylation of adenine in a GATC tetranucleotide which inhibits virulence of the bacteria. Thus, compounds which inhibit proliferation of bacteria are useful in treating bacterial infections.

CROSS-REFERENCE

[0001] This patent application is a continuation-in-part of U.S. patent application Ser. No. 09/612,116 filed Jul. 7, 2000 which is a continuation-in-part of U.S. patent application Ser. No. 09/495,614, filed Feb. 1, 2000, which claims the priority benefit of U.S. patent application Ser. Nos. 09/241,951, filed Feb. 2, 1999, converted to U.S. Provisional Ser. No. 60/183,043, and 09/305,603, filed May 5, 1999, converted to U.S. Provisional Ser. No. 60/198,250, all of which are incorporated by reference in their entirety and to which applications is claimed priority.

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Jun 20, 2002

PUBLICATION-DATE: June 20, 2002

INVENTOR-INFORMATION:

NAME	CITY	STATE	COUNTRY	RULE-47
Mahan, Michael J.	Santa Barbara	CA	US	
Heithoff, Douglas M.	Goleta	CA	US	
Low, David A.	Goleta	CA	US	
Sinsheimer, Robert L.	Santa Barbbra	CA	US	

APPL-NO: 09/ 927885 [PALM]
DATE FILED: August 9, 2001

RELATED-US-APPL-DATA:

Application 09/927885 is a continuation-in-part-of US application 09/612116, filed July 7, 2000, PENDING
Application 09/612116 is a continuation-in-part-of US application 09/495614, filed February 1, 2000, PENDING
Application is a non-provisional-of-provisional application 60/183043, filed February 2, 1999,
Application is a non-provisional-of-provisional application 60/198250, filed May 5, 1999,

INT-CL: [07] A61 K 31/00, A61 K 31/52

US-CL-PUBLISHED: 514/1; 514/263.4
US-CL-CURRENT: 514/1; 514/263.4

REPRESENTATIVE-FIGURES: NONE

ABSTRACT:

The virulence of bacterial strains and in particular pathogenic bacteria which infect human is reduced by an agent which alters the bacteria's native level or activity of DNA methyltransferase (Dam). The agent causes an alteration in the bacteria's native level of methylation of adenine in a GATC tetranucleotide which inhibits virulence of the bacteria. Thus, compounds and formulations thereof which reduce bacterial virulence inhibit proliferation of bacteria and are useful in treating bacterial infections, particularly in humans.

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L3: Entry 28 of 54

File: USPT

Oct 9, 2001

DOCUMENT-IDENTIFIER: US 6300084 B1

TITLE: Anti-mitotic agent screening process

CLAIMS:

1. A method for screening potential anti-mitotic agents comprising:

a) providing one or more purified proteins that are required for yeast mitotic spindle formation, wherein the one or more proteins are selected from the group consisting of a protein encoded by the yeast gene YGR113w (Dam I), and proteins provided in a complex of the Duo 1 protein encoded by the yeast gene YGL061 c and the Dam I protein encoded by the yeast gene YGR113w,

b) combining in vitro said one or more proteins with microtubules, and a potential anti-mitotic agent;

c) incubating said one or more proteins, said microtubules, and said anti-mitotic agent;

d) assaying the microtubules for binding of said one or more proteins; and

e) identifying an anti-mitotic agent, wherein said anti-mitotic agent inhibits binding of said one or more proteins to said microtubules.

2. The method according to claim 1, wherein said microtubules comprise bovine brain microtubules.

3. The method according to claim 1, wherein said microtubules comprise mammalian microtubules.

4. The method according to claim 1, wherein said assaying comprises pelleting said microtubules by ultracentrifugation followed by polyacrylamide gel electrophoresis of the pellets to assess the inhibition of binding of said one or more proteins to said microtubules.

5. The method according to claim 1, wherein said assaying comprises attaching a fluorescent probe to at least one of said one or more proteins and assessing the inhibition of binding of said one or more proteins to said microtubules by measuring the amount of said fluorescent probe on said microtubules.

6. The method according to claim 5, wherein measuring comprises Fluorescent Resonance Energy Transfer (FRET) or fluorescence anisotropy.

7. The method according to claim 1, wherein said one or more proteins required for mitotic spindle formation consists of the Dam1p protein encoded by the yeast gene YGR113w.

8. The method according to claim 1, wherein said one or more proteins required for mitotic spindle formation are provided in a complex of the Duo1 protein encoded by the yeast gene YGL161c and the Dam1p protein encoded by the yeast gene YGR113w.

9. The method according to claim 8, wherein the assay for binding is an assay for assessing the binding of said complex to said microtubules.

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L3: Entry 34 of 54

File: USPT

Jun 6, 2000

US-PAT-NO: 6072102

DOCUMENT-IDENTIFIER: US 6072102 A

TITLE: Reversible nuclear genetic system for male sterility in transgenic plants

DATE-ISSUED: June 6, 2000

INVENTOR-INFORMATION:

NAME	CITY	STATE	ZIP CODE	COUNTRY
Cigan; Andrew M.	Des Moines	IA		
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US-CL-CURRENT: [800/274](#); [536/24.1](#), [800/271](#), [800/275](#), [800/285](#), [800/288](#)

CLAIMS:

What is claimed is:

1. A recombinant DNA molecule comprising:

(i) a lexA DNA binding site embedded in a tissue specific promoter which is operatively linked to a first DNA sequence encoding a gene product which when expressed in a plant inhibits or disrupts pollen formation or function, and

(ii) a second DNA sequence encoding a lexA repressor, which DNA sequence is operatively linked to an inducible promoter.

2. The recombinant DNA molecule of claim 1, wherein the tissue specific promoter is an anther-specific promoter.

3. The recombinant DNA molecule of claim 1, wherein the gene product is a cytotoxin.

4. The recombinant DNA molecule of claim 1, wherein the gene product is a diphtheria toxin A-chain.

5. The recombinant DNA molecule of claim 1, wherein the first DNA sequence is a cell cycle division mutant gene.

6. The recombinant DNA molecule of claim 5, wherein the cell cycle division mutant gene is selected from the group consisting of CC gene from maize, WT gene and P68.

7. The recombinant DNA molecule of claim 1, wherein the gene product is a methylase.

8. The recombinant DNA molecule of claim 7, wherein the methylase is a DAM methylase.

9. The recombinant DNA molecule of claim 1, wherein the inducible promoter is inducible by a chemical herbicidal safener.

10. The recombinant DNA molecule of claim 1, wherein the tissue specific promoter is an anther-specific promoter, the gene product is a DAM methylase, and the inducible promoter is inducible by a chemical herbicidal safener.

11. A plant cell comprising the recombinant DNA molecule of claim 1.

12. A plant comprising the recombinant DNA molecule of claim 1.

13. A method for producing reversible male sterility in a plant, comprising:

(a) providing a plant comprising a recombinant DNA molecule comprising

(i) a lexA DNA binding site embedded in a tissue specific promoter which is operatively linked to a first DNA sequence encoding a gene product which when expressed in a plant inhibits or disrupts pollen formation or function, and

(ii) a second DNA sequence encoding a lexA repressor, which DNA sequence is operatively linked to an inducible promoter; and

(b) exposing the plant to an inducer to reverse the male sterile effect of the recombinant DNA molecule.

14. The method of claim 13, wherein the tissue specific promoter is an anther-specific promoter.

15. The method of claim 13, wherein the gene product is a cytotoxin.

16. The method of claim 13, wherein the gene product is a methylase.

17. The method of claim 16, wherein the methylase is a DAM methylase.

18. The method of claim 13, wherein the inducer is an chemical herbicidal safener.

19. The method of claim 13, wherein the tissue specific promoter is an anther-specific promoter, the gene product is a DAM methylase, and the chemical inducer is a chemical inducer.

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